

RAW SEQUENCE LISTING

**The Biotechnology Systems Branch of the Scientific and Technical
Information Center (STIC) no errors detected.**

Application Serial Number: 10/578,400
Source: IFWO
Date Processed by STIC: 05/17/2006

ENTERED

CRF Errors Edited by the STIC Systems Branch

Serial Number: 10/578,400

CRF Edit Date: DA
Edited by: 05/17/2006

___ Realigned nucleic acid/amino acid numbers/text in cases where the sequence text "wrapped" to the next line

___ Corrected the SEQ ID NO. Sequence numbers edited were:

___ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited:

/ Deleted: / invalid beginning/end-of-file text ; ___ page numbers

___ Inserted mandatory headings/numeric identifiers, specifically:

___ Moved responses to same line as heading/numeric identifier, specifically:

___ Other:



IFWO

RAW SEQUENCE LISTING

DATE: 05/17/2006

PATENT APPLICATION: US/10/578,400

TIME: 09:52:40

Input Set : A:\pto.da.txt

Output Set: N:\CRF4\03162006\J578400.raw

```

4 <110> APPLICANT: Long, Li
5     Luqman, Mohammad
6     Yabannavar, Asha
7     Zaror, Isabel
9 <120> TITLE OF INVENTION: Methods of Therapy for Solid Tumors
10    Expressing the CD40 Cell-Surface Antigen
13 <130> FILE REFERENCE: PP21566.002 (282914)
C--> 15 <140> CURRENT APPLICATION NUMBER: US/10/578,400
C--> 15 <141> CURRENT FILING DATE: 2006-05-03
15 <150> PRIOR APPLICATION NUMBER: 60/565,634
16 <151> PRIOR FILING DATE: 2004-04-27
18 <150> PRIOR APPLICATION NUMBER: 60/565,710
19 <151> PRIOR FILING DATE: 2004-04-27
21 <150> PRIOR APPLICATION NUMBER: 60/525,579
22 <151> PRIOR FILING DATE: 2003-11-26
24 <150> PRIOR APPLICATION NUMBER: 60/517,337
25 <151> PRIOR FILING DATE: 2003-11-04
27 <160> NUMBER OF SEQ ID NOS: 12
29 <170> SOFTWARE: FastSEQ for Windows Version 4.0
31 <210> SEQ ID NO: 1
32 <211> LENGTH: 720
33 <212> TYPE: DNA
34 <213> ORGANISM: Artificial Sequence
36 <220> FEATURE:
37 <223> OTHER INFORMATION: Coding sequence for light chain of 12.12 human
38    anti-CD40 antibody
W--> 41 <221> NAME/KEY: CDS
42 <222> LOCATION: (1)...(720)
W--> 44 <400> 1
45 atg gcg ctc cct gct cag ctc ctg ggg ctg cta atg ctc tgg gtc tct   48
46 Met Ala Leu Pro Ala Gln Leu Leu Gly Leu Leu Met Leu Trp Val Ser
47 1             5             10             15
49 gga tcc agt ggg gat att gtg atg act cag tct cca ctc tcc ctg acc   96
50 Gly Ser Ser Gly Asp Ile Val Met Thr Gln Ser Pro Leu Ser Leu Thr
51             20             25             30
53 gtc acc cct gga gag ccg gcc tcc atc tcc tgc agg tcc agt cag agc   144
54 Val Thr Pro Gly Glu Pro Ala Ser Ile Ser Cys Arg Ser Ser Gln Ser
55             35             40             45
57 ctc ctg tat agt aat gga tac aac tat ttg gat tgg tac ctg cag aag   192
58 Leu Leu Tyr Ser Asn Gly Tyr Asn Tyr Leu Asp Trp Tyr Leu Gln Lys
59             50             55             60
61 cca ggg cag tct cca cag gtc ctg atc tct ttg ggt tct aat cgg gcc   240
62 Pro Gly Gln Ser Pro Gln Val Leu Ile Ser Leu Gly Ser Asn Arg Ala

```

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Input Set : A:\pto.da.txt

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```

63 65          70          75          80
65 tcc ggg gtc cct gac agg ttc agt ggc agt gga tca ggc aca gat ttt 288
66 Ser Gly Val Pro Asp Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp Phe
67          85          90          95
69 aca ctg aaa atc agc aga gtg gag gct gag gat gtt ggg gtt tat tac 336
70 Thr Leu Lys Ile Ser Arg Val Glu Ala Glu Asp Val Gly Val Tyr Tyr
71          100          105          110
73 tgc atg caa gct cga caa act cca ttc act ttc ggc cct ggg acc aaa 384
74 Cys Met Gln Ala Arg Gln Thr Pro Phe Thr Phe Gly Pro Gly Thr Lys
75          115          120          125
77 gtg gat atc aga cga act gtg gct gca cca tct gtc ttc atc ttc ccg 432
78 Val Asp Ile Arg Arg Thr Val Ala Ala Pro Ser Val Phe Ile Phe Pro
79          130          135          140
81 cca tct gat gag cag ttg aaa tct gga act gcc tct gtt gtg tgc ctg 480
82 Pro Ser Asp Glu Gln Leu Lys Ser Gly Thr Ala Ser Val Val Cys Leu
83 145          150          155          160
85 ctg aat aac ttc tat ccc aga gag gcc aaa gta cag tgg aag gtg gat 528
86 Leu Asn Asn Phe Tyr Pro Arg Glu Ala Lys Val Gln Trp Lys Val Asp
87          165          170          175
89 aac gcc ctc caa tcg ggt aac tcc cag gag agt gtc aca gag cag gac 576
90 Asn Ala Leu Gln Ser Gly Asn Ser Gln Glu Ser Val Thr Glu Gln Asp
91          180          185          190
93 agc aag gac agc acc tac agc ctc agc agc acc ctg acg ctg agc aaa 624
94 Ser Lys Asp Ser Thr Tyr Ser Leu Ser Ser Thr Leu Thr Leu Ser Lys
95          195          200          205
97 gca gac tac gag aaa cac aaa gtc tac gcc tgc gaa gtc acc cat cag 672
98 Ala Asp Tyr Glu Lys His Lys Val Tyr Ala Cys Glu Val Thr His Gln
99          210          215          220
101 ggc ctg agc tcg ccc gtc aca aag agc ttc aac agg gga gag tgt tag 720
102 Gly Leu Ser Ser Pro Val Thr Lys Ser Phe Asn Arg Gly Glu Cys *
103 225          230          235
107 <210> SEQ ID NO: 2
108 <211> LENGTH: 239
109 <212> TYPE: PRT
110 <213> ORGANISM: Artificial Sequence
112 <220> FEATURE:
113 <223> OTHER INFORMATION: Light chain of 12.12 human anti-CD40 antibody
115 <400> SEQUENCE: 2
116 Met Ala Leu Pro Ala Gln Leu Leu Gly Leu Leu Met Leu Trp Val Ser
117 1          5          10          15
118 Gly Ser Ser Gly Asp Ile Val Met Thr Gln Ser Pro Leu Ser Leu Thr
119          20          25          30
120 Val Thr Pro Gly Glu Pro Ala Ser Ile Ser Cys Arg Ser Ser Gln Ser
121          35          40          45
122 Leu Leu Tyr Ser Asn Gly Tyr Asn Tyr Leu Asp Trp Tyr Leu Gln Lys
123          50          55          60
124 Pro Gly Gln Ser Pro Gln Val Leu Ile Ser Leu Gly Ser Asn Arg Ala
125 65          70          75          80
126 Ser Gly Val Pro Asp Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp Phe

```

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PATENT APPLICATION: US/10/578,400

TIME: 09:52:40

Input Set : A:\pto.da.txt

Output Set: N:\CRF4\05162006\J578400.raw

```

127      85      90      95
128 Thr Leu Lys Ile Ser Arg Val Glu Ala Glu Asp Val Gly Val Tyr Tyr
129      100      105      110
130 Cys Met Gln Ala Arg Gln Thr Pro Phe Thr Phe Gly Pro Gly Thr Lys
131      115      120      125
132 Val Asp Ile Arg Arg Thr Val Ala Ala Pro Ser Val Phe Ile Phe Pro
133      130      135      140
134 Pro Ser Asp Glu Gln Leu Lys Ser Gly Thr Ala Ser Val Val Cys Leu
135 145      150      155      160
136 Leu Asn Asn Phe Tyr Pro Arg Glu Ala Lys Val Gln Trp Lys Val Asp
137      165      170      175
138 Asn Ala Leu Gln Ser Gly Asn Ser Gln Glu Ser Val Thr Glu Gln Asp
139      180      185      190
140 Ser Lys Asp Ser Thr Tyr Ser Leu Ser Ser Thr Leu Thr Leu Ser Lys
141      195      200      205
142 Ala Asp Tyr Glu Lys His Lys Val Tyr Ala Cys Glu Val Thr His Gln
143      210      215      220
144 Gly Leu Ser Ser Pro Val Thr Lys Ser Phe Asn Arg Gly Glu Cys
145 225      230      235
148 <210> SEQ ID NO: 3
149 <211> LENGTH: 2016
150 <212> TYPE: DNA
151 <213> ORGANISM: Artificial Sequence
153 <220> FEATURE:
154 <223> OTHER INFORMATION: Coding sequence for heavy chain of 12.12 human
155      anti-CD40 antibody (with introns)
158 <400> SEQUENCE: 3
159 atggagtttg ggctgagctg ggttttcctt gttgctattt taagaggtgt ccagtgtcag 60
160 gtgcagttgg tggagtctgg gggaggcgtg gtccagcctg ggaggtcctt gagactctcc 120
161 tgtgcagcct ctggattcac cttcagtagc tatggcatgc actgggtccg ccaggctcca 180
162 ggcaaggggc tggagtgggt ggcagttata tcatatgagg aaagtaatag ataccatgca 240
163 gactccgtga agggccgatt caccatctcc agagacaatt ccaagatcac gctgtatctg 300
164 caaatgaaca gcctcagaac tgaggacacg gctgtgtatt actgtgcgag agatgggggt 360
165 atagcagcac ctgggcctga ctactggggc cagggaaccc tggtcaccgt ctctcagca 420
166 agtaccaagg gcccatccgt cttccccctg gcgcccgcta gcaagagcac ctctgggggc 480
167 acagcggccc tgggctgcct ggtcaaggac tacttccccg aaccggtgac ggtgtcgtgg 540
168 aactcaggcg cctgaccag cggcgtgcac accttcccgg ctgtcctaca gtcctcagga 600
169 ctctactccc tcagcagcgt ggtgaccgtg ccctccagca gcttggggcac ccagacctac 660
170 atctgcaacg tgaatcaciaa gcccagcaac accaagggtg acaagagagt tggtgagagg 720
171 ccagcacagg gagggagggt gtctgctgga agccaggctc agcgtctctg cctggacgca 780
172 tcccggtat gcagtccag tccagggcag caaggcaggc cccgtctgcc tcttcacccg 840
173 gaggcctctg cccgccccac tcatgctcag ggagagggtc ttctggcttt tttcccaggc 900
174 tctgggcagg cacaggctag gtgcccctaa cccaggccct gcacacaaag gggcagggtg 960
175 tgggctcaga cctgccaaga gccatatccg ggaggaccct gcccctgacc taagcccacc 1020
176 ccaaaggcca aactctccac tccctcagct cggacacctt ctctcctccc agattccagt 1080
177 aactcccaat cttctctctg cagagcccaa atcttgtgac aaaactcaca catgcccacc 1140
178 gtgcccagggt aagccagccc aggcctcgcc ctccagctca aggcgggaca ggtgccctag 1200
179 agtagcctgc atccagggac aggccccagc cgggtgctga cacgtccacc tccatctctt 1260
180 cctcagcacc tgaactcctg gggggaccgt cagtcttctt cttcccccca aaacccaagg 1320

```

RAW SEQUENCE LISTING

DATE: 05/17/2006

PATENT APPLICATION: US/10/578,400

TIME: 09:52:40

Input Set : A:\pto.da.txt

Output Set: N:\CRF4\05162006\J578400.raw

```

181 acaccctcat gatctcccg g acccctgagg tcacatgcgt ggtggtggac gtgagccacg 1380
182 aagaccctga ggtcaagttc aactggtacg tggacggcgt ggaggtgcat aatgccaaga 1440
183 caaagccgcg ggaggagcag tacaacagca cgtaccgtgt ggtcagcgtc ctcaccgtcc 1500
184 tgcaccagga ctggctgaat ggcaaggagt acaagtgcaa ggtctccaac aaagccctcc 1560
185 cagcccccac cgagaaaacc atctccaaag ccaaagggtg gacccgtggg gtgcgagggc 1620
186 cacatggaca gaggccggct cggccccacc tctgccctga gagtgaccgc tgtaccaacc 1680
187 tctgtcccta cagggcagcc ccgagaacca caggtgtaca ccctgcccc atccccggag 1740
188 gagatgacca agaaccaggt cagcctgacc tgccigtgca aaggcttcta tcccagcgac 1800
189 atcgccgtgg agtgggagag caatgggcag ccggagaaca actacaagac cagcctccc 1860
190 gtgctggact ccgacggctc cttcttcctc tatagcaagc tcaccgtgga caagagcagg 1920
191 tggcagcagg ggaacgtctt ctcattgctc gtgatgcatg aggctctgca caaccactac 1980
192 acgcagaaga gcctctccct gtctccgggt aatga 2016

```

194 <210> SEQ ID NO: 4

195 <211> LENGTH: 469

196 <212> TYPE: PRT

197 <213> ORGANISM: Artificial Sequence

199 <220> FEATURE:

200 <223> OTHER INFORMATION: Heavy chain of 12.12 human anti-CD40 antibody

202 <400> SEQUENCE: 4

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203 Met Glu Phe Gly Leu Ser Trp Val Phe Leu Val Ala Ile Leu Arg Gly
204 1 5 10 15
205 Val Gln Cys Gln Val Gln Leu Val Glu Ser Gly Gly Gly Val Val Gln
206 20 25 30
207 Pro Gly Arg Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe
208 35 40 45
209 Ser Ser Tyr Gly Met His Trp Val Arg Gln Ala Pro Gly Lys Gly Leu
210 50 55 60
211 Glu Trp Val Ala Val Ile Ser Tyr Glu Glu Ser Asn Arg Tyr His Ala
212 65 70 75 80
213 Asp Ser Val Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Ile
214 85 90 95
215 Thr Leu Tyr Leu Gln Met Asn Ser Leu Arg Thr Glu Asp Thr Ala Val
216 100 105 110
217 Tyr Tyr Cys Ala Arg Asp Gly Gly Ile Ala Ala Pro Gly Pro Asp Tyr
218 115 120 125
219 Trp Gly Gln Gly Thr Leu Val Thr Val Ser Ser Ala Ser Thr Lys Gly
220 130 135 140
221 Pro Ser Val Phe Pro Leu Ala Pro Ala Ser Lys Ser Thr Ser Gly Gly
222 145 150 155 160
223 Thr Ala Ala Leu Gly Cys Leu Val Lys Asp Tyr Phe Pro Glu Pro Val
224 165 170 175
225 Thr Val Ser Trp Asn Ser Gly Ala Leu Thr Ser Gly Val His Thr Phe
226 180 185 190
227 Pro Ala Val Leu Gln Ser Ser Gly Leu Tyr Ser Leu Ser Ser Val Val
228 195 200 205
229 Thr Val Pro Ser Ser Ser Leu Gly Thr Gln Thr Tyr Ile Cys Asn Val
230 210 215 220
231 Asn His Lys Pro Ser Asn Thr Lys Val Asp Lys Arg Val Glu Pro Lys
232 225 230 235 240

```

RAW SEQUENCE LISTING

DATE: 05/17/2006

PATENT APPLICATION: US/10/578,400

TIME: 09:52:40

Input Set : A:\pto.da.txt

Output Set: N:\CRF4\05162006\J578400.raw

```

233 Ser Cys Asp Lys Thr His Thr Cys Pro Pro Cys Pro Ala Pro Glu Leu
234                245                250                255
235 Leu Gly Gly Pro Ser Val Phe Leu Phe Pro Pro Lys Pro Lys Asp Thr
236                260                265                270
237 Leu Met Ile Ser Arg Thr Pro Glu Val Thr Cys Val Val Val Asp Val
238                275                280                285
239 Ser His Glu Asp Pro Glu Val Lys Phe Asn Trp Tyr Val Asp Gly Val
240                290                295                300
241 Glu Val His Asn Ala Lys Thr Lys Pro Arg Glu Glu Gln Tyr Asn Ser
242 305                310                315                320
243 Thr Tyr Arg Val Val Ser Val Leu Thr Val Leu His Gln Asp Trp Leu
244                325                330                335
245 Asn Gly Lys Glu Tyr Lys Cys Lys Val Ser Asn Lys Ala Leu Pro Ala
246                340                345                350
247 Pro Ile Glu Lys Thr Ile Ser Lys Ala Lys Gly Gln Pro Arg Glu Pro
248                355                360                365
249 Gln Val Tyr Thr Leu Pro Pro Ser Arg Glu Glu Met Thr Lys Asn Gln
250                370                375                380
251 Val Ser Leu Thr Cys Leu Val Lys Gly Phe Tyr Pro Ser Asp Ile Ala
252 385                390                395                400
253 Val Glu Trp Glu Ser Asn Gly Gln Pro Glu Asn Asn Tyr Lys Thr Thr
254                405                410                415
255 Pro Pro Val Leu Asp Ser Asp Gly Ser Phe Phe Leu Tyr Ser Lys Leu
256                420                425                430
257 Thr Val Asp Lys Ser Arg Trp Gln Gln Gly Asn Val Phe Ser Cys Ser
258                435                440                445
259 Val Met His Glu Ala Leu His Asn His Tyr Thr Gln Lys Ser Leu Ser
260                450                455                460
261 Leu Ser Pro Gly Lys
262 465
265 <210> SEQ ID NO: 5
266 <211> LENGTH: 469
267 <212> TYPE: PRT
268 <213> ORGANISM: Artificial Sequence
270 <220> FEATURE:
271 <223> OTHER INFORMATION: Heavy chain of variant of 12.12 human anti-CD40
272 antibody
274 <400> SEQUENCE: 5
275 Met Glu Phe Gly Leu Ser Trp Val Phe Leu Val Ala Ile Leu Arg Gly
276 1                5                10                15
277 Val Gln Cys Gln Val Gln Leu Val Glu Ser Gly Gly Gly Val Val Gln
278                20                25                30
279 Pro Gly Arg Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe
280                35                40                45
281 Ser Ser Tyr Gly Met His Trp Val Arg Gln Ala Pro Gly Lys Gly Leu
282                50                55                60
283 Glu Trp Val Ala Val Ile Ser Tyr Glu Glu Ser Asn Arg Tyr His Ala
284 65                70                75                80
285 Asp Ser Val Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Ile

```

VERIFICATION SUMMARY

DATE: 05/17/2006

PATENT APPLICATION: US/10/578,400

TIME: 09:52:41

Input Set : A:\pto.da.txt

Output Set: N:\CRF4\05162006\J578400.raw

L:15 M:270 C: Current Application Number differs, Replaced Current Application No
L:15 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:41 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:44 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:1
L:537 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:9
L:638 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:11

STATISTICS SUMMARY

PATENT APPLICATION: US/10/578,400

DATE: 05/17/2006

TIME: 09:52:41

Input Set : A:\pto.da.txt

Output Set: N:\CRF4\05162006\J578400.raw

Application Serial Number: US/10/578,400

Alpha or Numeric or Xml: Numeric

Application Class:

Application File Date: 05-03-2006

Art Unit: IFWO

Software Application: FastSEQ

Total Number of Sequences: 12

Total Nucleotides: 4132

Total Amino Acids: 2844

Number of Errors: 0

Number of Warnings: 4

Number of Corrections: 2

MESSAGE SUMMARY

258 W: 3 (Mandatory Feature missing)

270 C: 1 (Current Application Number differs)

271 C: 1 (Current Filing Date differs)

281 W: 1 (Numeric Fields not Ordered)

Raw Sequence Listing before editing,
for reference only



IFWP

RAW SEQUENCE LISTING

DATE: 05/15/2006

PATENT APPLICATION: US/10/578,400

TIME: 09:47:23

Input Set : A:\sequence lstg.txt

Output Set: N:\CRF4\05152006\J578400.raw

4 <110> APPLICANT: Long, Li
 5 Luqman, Mohammad
 6 Yabannavar, Asha
 7 Zaror, Isabel
 9 <120> TITLE OF INVENTION: Methods of Therapy for Solid Tumors
 10 Expressing the CD40 Cell-Surface Antigen
 13 <130> FILE REFERENCE: PP21566.002 (282914)
 C--> 15 <140> CURRENT APPLICATION NUMBER: US/10/578,400
 C--> 15 <141> CURRENT FILING DATE: 2006-05-03
 15 <150> PRIOR APPLICATION NUMBER: 60/565,634
 16 <151> PRIOR FILING DATE: 2004-04-27
 18 <150> PRIOR APPLICATION NUMBER: 60/565,710
 19 <151> PRIOR FILING DATE: 2004-04-27
 21 <150> PRIOR APPLICATION NUMBER: 60/525,579
 22 <151> PRIOR FILING DATE: 2003-11-26
 24 <150> PRIOR APPLICATION NUMBER: 60/517,337
 25 <151> PRIOR FILING DATE: 2003-11-04
 27 <160> NUMBER OF SEQ ID NOS: 12
 29 <170> SOFTWARE: FastSEQ for Windows Version 4.0

Does Not Comply
 Corrected Diskette Needed
 CP9-2)

ERRORED SEQUENCES

712 <210> SEQ ID NO: 12
 713 <211> LENGTH: 277
 714 <212> TYPE: PRT
 715 <213> ORGANISM: Homo sapiens
 717 <400> SEQUENCE: 12
 718 Met Val Arg Leu Pro Leu Gln Cys Val Leu Trp Gly Cys Leu Leu Thr
 719 1 5 10 15
 720 Ala Val His Pro Glu Pro Pro Thr Ala Cys Arg Glu Lys Gln Tyr Leu
 721 20 25 30
 722 Ile Asn Ser Gln Cys Cys Ser Leu Cys Gln Pro Gly Gln Lys Leu Val
 723 35 40 45
 724 Ser Asp Cys Thr Glu Phe Thr Glu Thr Glu Cys Leu Pro Cys Gly Glu
 725 50 55 60
 726 Ser Glu Phe Leu Asp Thr Trp Asn Arg Glu Thr His Cys His Gln His
 727 65 70 75 80
 728 Lys Tyr Cys Asp Pro Asn Leu Gly Leu Arg Val Gln Gln Lys Gly Thr
 729 85 90 95
 730 Ser Glu Thr Asp Thr Ile Cys Thr Cys Glu Glu Gly Trp His Cys Thr
 731 100 105 110
 732 Ser Glu Ala Cys Glu Ser Cys Val Leu His Arg Ser Cys Ser Pro Gly

RAW SEQUENCE LISTING

DATE: 05/15/2006

PATENT APPLICATION: US/10/578,400

TIME: 09:47:23

Input Set : A:\sequence 1stg.txt

Output Set: N:\CRF4\05152006\J578400.raw

733 115 120 125
734 Phe Gly Val Lys Gln Ile Ala Thr Gly Val Ser Asp Thr Ile Cys Glu
735 130 135 140
736 Pro Cys Pro Val Gly Phe Phe Ser Asn Val Ser Ser Ala Phe Glu Lys
737 145 150 155 160
738 Cys His Pro Trp Thr Ser Cys Glu Thr Lys Asp Leu Val Val Gln Gln
739 165 170 175
740 Ala Gly Thr Asn Lys Thr Asp Val Val Cys Gly Pro Gln Asp Arg Leu
741 180 185 190
742 Arg Ala Leu Val Val Ile Pro Ile Ile Phe Gly Ile Leu Phe Ala Ile
743 195 200 205
744 Leu Leu Val Leu Val Phe Ile Lys Lys Val Ala Lys Lys Pro Thr Asn
745 210 215 220
746 Lys Ala Pro His Pro Lys Gln Glu Pro Gln Glu Ile Asn Phe Pro Asp
747 225 230 235 240
748 Asp Leu Pro Gly Ser Asn Thr Ala Ala Pro Val Gln Glu Thr Leu His
749 245 250 255
750 Gly Cys Gln Pro Val Thr Gln Glu Asp Gly Lys Glu Ser Arg Ile Ser
751 260 265 270
752 Val Gln Glu Arg Gln
753 275
E--> 755 12
761 RTA01/2168201v1

deleted

VERIFICATION SUMMARY

DATE: 05/15/2006

PATENT APPLICATION: US/10/578,400

TIME: 09:47:24

Input Set : A:\sequence 1stg.txt

Output Set: N:\CRF4\05152006\J578400.raw

L:15 M:270 C: Current Application Number differs, Replaced Current Application No
L:15 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:41 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:44 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:1
L:537 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:9
L:638 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:11
L:755 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:12